

## SEQUENCE LISTING

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<120> SUPERFICIAL ZONE PROTEIN AND METHODS OF  
 MAKING AND USING SAME

<130> 07083.0008U5

<150> 60/258,920

<151> 2000-12-29

<160> 11

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =  
 synthetic construct

<400> 1

Asp Glu Ala Gly Ser Gly  
 1 5

<210> 2

<211> 188

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =  
 synthetic construct

<400> 2

Met	Ala	Trp	Lys	Thr	Leu	Pro	Ile	Tyr	Leu	Leu	Leu	Leu	Leu	Ser	Val
1				5				10						15	
Phe	Val	Ile	Gln	Gln	Val	Ser	Ser	Gln	Asp	Leu	Ser	Ser	Cys	Ala	Gly
		20						25					30		
Arg	Cys	Gly	Glu	Gly	Tyr	Ser	Arg	Asp	Ala	Thr	Cys	Asn	Cys	Asp	Tyr
		35					40					45			

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Asn Cys Gln His Tyr Met Glu Cys Cys Pro Asp Phe Lys Arg Val Cys  
 50 55 60  
 Thr Ala Glu Leu Ser Cys Lys Gly Arg Cys Phe Glu Ser Phe Glu Arg  
 65 70 75 80  
 Gly Arg Glu Cys Asp Cys Asp Ala Gln Cys Lys Lys Tyr Asp Lys Cys  
 85 90 95  
 Cys Pro Asp Tyr Glu Ser Phe Cys Ala Glu Val Lys Asp Asn Lys Lys  
 100 105 110  
 Asn Arg Thr Lys Lys Lys Pro Thr Pro Lys Pro Pro Val Val Asp Glu  
 115 120 125  
 Ala Gly Ser Gly Leu Asp Asn Gly Asp Phe Lys Val Thr Thr Pro Asp  
 130 135 140  
 Thr Ser Thr Thr Gln His Asn Lys Val Ser Thr Ser Pro Lys Ile Thr  
 145 150 155 160  
 Thr Ala Lys Pro Ile Asn Pro Arg Pro Gln Ser Ser Pro Asn Ser Asp  
 165 170 175  
 Thr Ser Lys Glu Thr Ser Leu Thr Val Asn Lys Glu  
 180 185

<210> 3

<211> 538

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =  
synthetic construct

<400> 3

Pro Thr Thr Ile His Lys Ser Pro Asp Glu Ser Thr Pro Glu Leu Ser  
 1 5 10 15  
 Ala Glu Pro Thr Pro Lys Ala Leu Glu Asn Ser Pro Lys Glu Pro Gly  
 20 25 30  
 Val Pro Thr Thr Lys Thr Pro Ala Ala Thr Lys Pro Glu Met Thr Thr  
 35 40 45  
 Thr Ala Lys Asp Lys Thr Thr Glu Arg Asp Leu Arg Thr Thr Pro Glu  
 50 55 60  
 Thr Thr Thr Ala Ala Pro Lys Met Thr Lys Glu Thr Ala Thr Thr Thr  
 65 70 75 80  
 Glu Lys Thr Thr Glu Ser Lys Ile Thr Ala Thr Thr Thr Gln Val Thr  
 85 90 95  
 Ser Thr Thr Thr Gln Asp Thr Thr Pro Phe Lys Ile Thr Thr Leu Lys  
 100 105 110  
 Thr Thr Leu Ala Pro Lys Val Thr Thr Thr Lys Lys Thr Ile Thr Thr  
 115 120 125  
 Thr Glu Ile Met Asn Lys Pro Glu Glu Thr Ala Lys Pro Lys Asp Arg  
 130 135 140  
 Ala Thr Asn Ser Lys Ala Thr Thr Pro Lys Pro Gln Lys Pro Thr Lys  
 145 150 155 160  
 Ala Pro Lys Lys Pro Thr Ser Thr Lys Lys Pro Lys Thr Met Pro Arg  
 165 170 175  
 Val Arg Lys Pro Lys Thr Thr Pro Thr Pro Arg Lys Met Thr Ser Thr  
 180 185 190

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Met Pro Glu Leu Asn Pro Thr Ser Arg Ile Ala Glu Ala Met Leu Gln
      195                      200                      205
Thr Thr Thr Arg Pro Asn Gln Thr Pro Asn Ser Lys Leu Val Glu Val
      210                      215                      220
Asn Pro Lys Ser Glu Asp Ala Gly Gly Ala Glu Gly Glu Thr Pro His
225                      230                      235                      240
Met Leu Leu Arg Pro His Val Phe Met Pro Glu Val Thr Pro Asp Met
      245                      250                      255
Asp Tyr Leu Pro Arg Val Pro Asn Gln Gly Ile Ile Ile Asn Pro Met
      260                      265                      270
Leu Ser Asp Glu Thr Asn Ile Cys Asn Gly Lys Pro Val Asp Gly Leu
      275                      280                      285
Thr Thr Leu Arg Asn Gly Thr Leu Val Ala Phe Arg Gly His Tyr Phe
      290                      295                      300
Trp Met Leu Ser Pro Phe Ser Pro Pro Ser Pro Ala Arg Arg Ile Thr
305                      310                      315                      320
Glu Val Trp Gly Ile Pro Ser Pro Ile Asp Thr Val Phe Thr Arg Cys
      325                      330                      335
Asn Cys Glu Gly Lys Thr Phe Phe Phe Lys Asp Ser Gln Tyr Trp Arg
      340                      345                      350
Phe Thr Asn Asp Ile Lys Asp Ala Gly Tyr Pro Lys Pro Ile Phe Lys
      355                      360                      365
Gly Phe Gly Gly Leu Thr Gly Gln Ile Val Ala Ala Leu Ser Thr Ala
      370                      375                      380
Lys Tyr Lys Asn Trp Pro Glu Ser Val Tyr Phe Phe Lys Arg Gly Gly
385                      390                      395                      400
Ser Ile Gln Gln Tyr Ile Tyr Lys Gln Glu Pro Val Gln Lys Cys Pro
      405                      410                      415
Gly Arg Arg Pro Ala Leu Asn Tyr Pro Val Tyr Gly Glu Met Thr Gln
      420                      425                      430
Val Arg Arg Arg Arg Phe Glu Arg Ala Ile Gly Pro Ser Gln Thr His
      435                      440                      445
Thr Ile Arg Ile Gln Tyr Ser Pro Ala Arg Leu Ala Tyr Gln Asp Lys
      450                      455                      460
Gly Val Leu His Asn Glu Val Lys Val Ser Ile Leu Trp Arg Gly Leu
465                      470                      475                      480
Pro Asn Val Val Thr Ser Ala Ile Ser Leu Pro Asn Ile Arg Lys Pro
      485                      490                      495
Asp Gly Tyr Asp Tyr Tyr Ala Phe Ser Lys Asp Gln Tyr Tyr Asn Ile
      500                      505                      510
Asp Val Pro Ser Arg Thr Ala Arg Ala Ile Thr Thr Arg Ser Gly Gln
      515                      520                      525
Thr Leu Ser Lys Val Trp Tyr Asn Cys Pro
      530                      535

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&lt;210&gt; 4

&lt;211&gt; 3

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

<223> Description of Artificial Sequence; note =  
synthetic construct

10033694.23101

<221> VARIANT

<222> 2

<223> Xaa is any amino acid except Pro

<221> VARIANT

<222> 3

<223> Xaa is either Thr or Ser

<400> 4

Asn Xaa Xaa

1

<210> 5

<211> 488

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =  
synthetic construct

<400> 5

atgcatgaaa	cattccatta	ctgtgttctg	tgttgtttct	gatcacaatt	tatccaaatt	60
atcagcgtga	ggagagtggg	agggatttag	gatccactga	acgtgttaaa	cgtcacatac	120
tgggtgtgcc	tgtttaagga	gctgactcgg	gcttcgtaa	ggcgcgcttg	atcctcggag	180
gggggggtgg	acgcgcgcca	agtagaatat	acagtgtgtc	cgttagaggt	ttctgtgcag	240
aagtaaaaga	taacaagaag	aacagaacta	aaaagaaacc	tacccccaaa	ccaccagttg	300
tagatgaagc	tggaagtgga	ttggacaatg	gtgacttcaa	ggtcacaact	cctgacacgt	360
ctaccacca	acacaataaa	gtcagcacat	ctcccaagat	cacaacagca	aaaccaataa	420
atcccagacc	ccagtcttca	cctaattctg	atacatctaa	agagacgtct	ttgacagtga	480
ataaagag						488

<210> 6

<211> 1620

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =  
synthetic construct

<400> 6

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ccaaaagctc	ttgaaaacag	tcccaaggaa	cctgggtgtac	ctacaactaa	gactcctgca	120
gcgactaaac	ctgaaatgac	tacaacagct	aaagacaaga	caacagaaag	agacttacgt	180
actacacctg	aaactacaac	tgctgcacct	aagatgacaa	aagagacagc	aactacaaca	240
gaaaaaacta	ccgaatccaa	aataacagct	acaaccacac	aagtaacatc	taccacaact	300
caagatacca	caccattcaa	aattactact	cttaaaaacaa	ctactcttgc	acccaaaagta	360
actacaacaa	aaaagacaat	tactaccact	gagattatga	acaaacctga	agaaacagct	420
aaacccaaaag	acagagctac	taattctaaa	gcgacaactc	ctaaacctca	aaagccaacc	480

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aaagcaccca	aaaaaccac	ttctacaaa	aagccaaaaa	caatgcctag	agtgagaaaa	540
ccaaagacga	caccaactcc	ccgcaagatg	acatcaacaa	tgccagaatt	gaaccctacc	600
tcaagaatag	cagaagccat	gctccaaacc	accaccagac	ctaaccaaac	tccaaactcc	660
aaactagttg	aagtaaatac	aaagagtga	gatgcaggtg	gtgctgaagg	agaaacacct	720
catatgcttc	tcaggcccca	tgtgttcctg	cctgaagtta	ctcccgacat	ggattactta	780
ccgagagtac	ccaatcaagg	cattatcatc	aatcccatgc	tttccgatga	gaccaatata	840
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&lt;210&gt; 7

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

<223> Description of Artificial Sequence; note =  
synthetic construct

&lt;400&gt; 7

atggcatgga aaacacttcc catt

24

&lt;210&gt; 8

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

<223> Description of Artificial Sequence; note =  
synthetic construct

&lt;400&gt; 8

ctaaggacag ttgtaccaga cttt

24

&lt;210&gt; 9

&lt;211&gt; 4

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

<223> Description of Artificial Sequence; note =  
synthetic construct

10033694.123101

<400> 9  
Phe Ala Cys Glu  
1

<210> 10  
<211> 8  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence; note =  
synthetic construct

<400> 10  
Val Lys Asp Asn Lys Lys Asn Arg  
1 5

<210> 11  
<211> 8  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence; note =  
synthetic construct

<221> VARIANT  
<222> 8  
<223> Xaa is either Thr or Pro

<400> 11  
Lys Glu Pro Ala Pro Thr Thr Xaa  
1 5  
2  
5

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